RAW SEQUENCE LISTING PATENT APPLICATION US/09/506,079

DATE: 03/09/2000 TIME: 03:28:58

INPUT SET: S34978.raw

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

Does Not Comply
Corrected Diskette Needed

1 SEQUENCE LISTING seepp2,4 2 3 (1) General Information: 4 (i) APPLICANTS: Clinton, Gail M., Doherty, Joni Kristin, and Adelman, John P. 5 6 7 (ii) TITLE OF INVENTION: HER-2 BINDING ANTAGONISTS 8 9 (iii) NUMBER OF SEQUENCES: 10 10 (iv) CORRESPONDENCE ADDRESS: 11 12 (A) ADDRESSEE: DAVIS WRIGHT TREMAINE 13 (B) STREET: 1501 Fourth Avenue, 2600 Century Square 14 (C) CITY: Seattle 15 (D) STATE: Washington 16 (E) COUNTRY: U.S.A. 17 (F) ZIP: 98101 18 19 (v) COMPUTER READABLE FORM: (A) MEDIUM TYPE: Floppy disk 20 (B) COMPUTER: PC compatible 21 (C) OPERATING SYSTEM: Windows95 22 (D) SOFTWARE: Word 23 24 (vi) CURRENT APPLICATION DATA: 25 --> oK 26 (A) APPLICATION NUMBER: to be assigned (B) FILING DATE: 19 January 2000 27 28 (C) CLASSIFICATION: 29 30 (viii) ATTORNEY/AGENT INFORMATION: 31 (A) NAME: Oster, Jeffrey B. (B) REGISTRATION NUMBER: 32,585 32 33 (C) REFERENCE/DOCKET NUMBER: 49321-1 34 (ix) TELECOMMUNICATION INFORMATION: 35 36 (A) TELEPHONE: 206 628 7711 37 (B) TELEFAX: 206 628 7699 38

ERRORED SEQUENCES FOLLOW:

60 (2) INFORMATION FOR SEQ ID NO:2:

RAW SEQUENCE LISTING PATENT APPLICATION US/09/506,079

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INPUT SET: S34978.raw

```
(i) SEQUENCE CHARACTERISTICS:
 62
 63
                (A) LENGTH: 419
                (B) TYPE: amino acid
 64
 65
                (C) STRANDEDNESS: single
                (D) TOPOLOGY: unknown
 66
 67
          (ii) MOLECULE TYPE: polypeptide
 68
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
 69
     Met Glu Leu Ala Ala Leu Cys Arg Trp Gly Leu Leu Leu Ala Leu Leu
 70
 71
                                            10
 72
      Pro Pro Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr Asp Cys Lys
 73
                      20
                                           25
 74
     Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met Leu Arg His
 75
                                       40
 76
     Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu Leu Thr Tyr
 77
                                  55
 78
      Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile Gln Glu Val
 79
      Gln Gly Tyr Val Leu Cys Ala His Asn Gln Val Arg Gln Val Pro Leu
 80
 81
                          85
      Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu Asp Asn Tyr
 82
 83
                                           105
     Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Agr Agn Thr Thr Pro
 84
 85
                                       120
                  115
      Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln Leu Arg Ser
 86
 87
                                   135
 88
     Leu Thr Glu Cys Leu Lys Gly Gly Val Leu Ile Gln Arg Asn Pro Gln
 89
                              150
 90
     Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe His Lys Asn
 91
                          165
                                               170
     Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser Arg Ala Cys
 92
 93
                      180
                                           185
 94
     His Pro Cys Ser Pro Cys Cys Lys Gly Ser Arg Cys Trp Gly Glu Ser
95
                                      200
 96
     Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala Gly Gly Cys
 97
                                  215
98
     Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His Glu Gln Cys
99
                              230
100
     Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu Ala Cys Leu
101
                          245
                                               250
     His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro Ala Leu Val
102
103
                      260
                                           265
104
     Thr Tyr Asn Thr Asp Thr Phe Glu Ser Cys Pro Asn Pro Glu Gly Arg
105
                                      280
106
     Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr Asn Lys Leu
107
                                  295
     Ser Thr Asp Val Gly Ser Cys Thr Leu Val Cys Pro Leu His Asn Gln
108
109
                              310
                                                   315
     Glu Val Thr Ala Glu Asp Gly Thr Gln Arg Cys Glu Lys Cys Ser Lys
110
111
                          325
                                               330
112
     Pro Cys Ala Arg Val Gly Xaa His Ser Xaa Xaa Pro Arg Pro Ala Ala
113
```

Best Available Copy

PAGE: 3

RAW SEQUENCE LISTING PATENT APPLICATION US/09/506,079

DATE: 03/09/2000 TIME: 03:28:59

INPUT SET: S34978.raw	INPIIT	SET:	S34978	raw
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															12 0 2	DELI DO IVI OII WII
114	Val	Pro	Val	Pro	Xaa	Arg	Xaa	Gln	Pro	Xaa	Pro	Ala	His	Pro	Val	Leu
115				355					360					365		
116	Ser	Phe	Leu	Arg	Pro	Ser	Trp	Asp	Xaa	Val	Ser	Ala	Phe	Tyr	Ser	Leu
117			370					375					380			
118	Pro	Leu	Ala	Pro	Leu	Asp	Pro	Thr	Ser	Val	Xaa	Ile	Ser	Pro	Val	Ser
119		385					390					395				
120	Val	Gly	Arg	Gly	Xaa	Asp	Pro	Asp	Ala	His	Val	Ala	Val	Xaa	Leu	Ser
121	400					405					410					415
122	Arg	Tyr	Glu	Gly												
123													•			

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

		 			AGG Arg				 45
					GCC Ala				90
					TCT Ser				135
					GTC Val				180
					GCT Ala				225
	TAT Tvr		TGA						240

(25) delete at end of file

PAGE: 1

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/506,079

DATE: 03/09/2000 TIME: 03:28:59

INPUT SET: S34978.raw

Line	Error	Original Text
26	Wrong application Serial Number	(A) APPLICATION NUMBER: to be assigned
63	Entered (419) and Calc. Seq. Length (418) differ	(A) LENGTH: 419
84	Wrong Amino Acid Designator	Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Agn Agn T
84	Wrong Amino Acid Designator	Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Agn Agn T